

sequence listing.txt  
SEQUENCE LISTING

<110> The University of York

<120> Expansin Polypeptides

<130> P102868WO

<140> PCT/GB04/004058

<141> 2004-09-23

<150> 0322317.9

<151> 2003-09-24

<160> 24

<170> PatentIn version 3.1

<210> 1

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<212> DNA

<213> Craterostigma plantagineum

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agctgcggca gctgctttga aattaaatgc gcgagcagca ttagcggcg cggaatgg	300
tgcttgcgg gcggcagcat taccgtgacc gcgaccaact tttgcccgcc gaacaacgcg	360
ctgccgaaca acgcggggcg ctggtgcaac ccgccgctgc agcattttga tctgagccag	420
ccggtgtttc agcatattgc gcagtatcgc gcgggcattg tgccggtgag ctatcgccgc	480
gtgccgtgcc gccgccgcgg cggcattcgc tttaccatta acggccatag ctattttaac	540
ctggtgctga ttaccaacgt gggcgggcg ggcgatgtgc atgcgggtgag cattaaaggc	600
gcgaccaccg attggcagcc gatgagccgc aactggggcc agaactggca gagcaacgcg	660
aacccgaacg gccagcgcct gagcttttaa gtgaccacca gcgatggccg caccctggtg	720
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<212> PRT

<213> Craterostigma plantagineum

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sequence listing.txt

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20 25 30

Trp Thr Asp Ala His Ala Thr Phe Tyr Gly Gly Ser Asp Ala Ser Gly  
35 40 45

Thr Met Gly Gly Ala Cys Gly Tyr Gly Asn Leu Tyr Ser Gln Gly Tyr  
50 55 60

Gly Thr Asn Thr Ala Ala Leu Ser Thr Ala Leu Phe Asn Asn Gly Leu  
65 70 75 80

Ser Cys Gly Ser Cys Phe Glu Ile Lys Cys Ala Ser Ser Ile Ser Gly  
85 90 95

Gly Gly Lys Trp Cys Leu Pro Gly Gly Ser Ile Thr Val Thr Ala Thr  
100 105 110

Asn Phe Cys Pro Pro Asn Asn Ala Leu Pro Asn Asn Ala Gly Gly Trp  
115 120 125

Cys Asn Pro Pro Leu Gln His Phe Asp Leu Ser Gln Pro Val Phe Gln  
130 135 140

His Ile Ala Gln Tyr Arg Ala Gly Ile Val Pro Val Ser Tyr Arg Arg  
145 150 155 160

Val Pro Cys Arg Arg Arg Gly Gly Ile Arg Phe Thr Ile Asn Gly His  
165 170 175

Ser Tyr Phe Asn Leu Val Leu Ile Thr Asn Val Gly Gly Ala Gly Asp  
180 185 190

Val His Ala Val Ser Ile Lys Gly Ala Thr Thr Asp Trp Gln Pro Met  
195 200 205

Ser Arg Asn Trp Gly Gln Asn Gln Ser Asn Ala Asn Pro Asn Gly Gln  
210 215 220

Arg Leu Ser Phe Lys Val Thr Thr Ser Asp Gly Arg Thr Leu Val Ser  
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sequence listing.txt

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aacggcctga cctgcggcgc gtgctatgaa ctgacctgca acaacgatcc gcgcggctgg 180  
tgcctgagcg gcaccattat ggtgaccgcg accaactttt gcccgccgaa cccgagcctg 240  
ccgaacgata acggcggctg gtgcaacccg ccgcgccagc attttgatct ggcggaaccg 300  
gcgttttctgc agattgcgca gtataaagcg ggcattgtgc cgggtgaacta tcgccgcgtg 360  
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gtgctggtga ccaacgtggg cggcgtgggc gatgtgcata gcgtgagcat taaaggcagc 480  
aacggcggct ggcagccgct gagccgcaac tggggccaga actggcagag caacagctat 540  
ctgaacggcc agagcctgag ctttcaggtg accaccagcg atggccgcac cgtgaccagc 600  
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20 25 30  
Ala Leu Ser Thr Ala Leu Phe Asn Asn Gly Leu Thr Cys Gly Ala Cys  
35 40 45  
Tyr Glu Leu Thr Cys Asn Asn Asp Pro Arg Gly Trp Cys Leu Ser Gly  
50 55 60  
Thr Ile Met Val Thr Ala Thr Asn Phe Cys Pro Pro Asn Pro Ser Leu  
65 70 75 80  
Pro Asn Asp Asn Gly Gly Trp Cys Asn Pro Pro Arg Gln His Phe Asp  
85 90 95  
Leu Ala Glu Pro Ala Phe Leu Gln Ile Ala Gln Tyr Lys Ala Gly Ile  
Page 3

100

Val Pro Val Asn Tyr Arg Arg Val Pro Cys Gln Lys Lys Gly Gly Ile  
115 120 125

Arg Phe Thr Ile Asn Gly His Ser Phe Phe Asn Leu Val Leu Val Thr  
130 135 140

Asn Val Gly Gly Val Gly Asp Val His Ser Val Ser Ile Lys Gly Ser  
145 150 155 160

Asn Gly Gly Trp Gln Pro Leu Ser Arg Asn Trp Gly Gln Asn Trp Gln  
165 170 175

Ser Asn Ser Tyr Leu Asn Gly Gln Ser Leu Ser Phe Gln Val Thr Thr  
180 185 190

Ser Asp Gly Arg Thr Val Thr Ser Tyr Asp Val Ala Pro Arg Gly Trp  
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ttaacaacg gcctggcgtg cggcagctgc tatcaggtgc gctgcgaagg cggcccgaaa 180  
tggtgcgtgc gcggcggcga tcgcattatt accgtgaccg cgaccaactt ttgcccgcgc 240  
aactatgcgc tggcgaacga taacggcggc tggtgcaacc cgccgcgcca gcattttgat 300  
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tatcgccgcg tgagctgcgt gaaaaaaggc ggcattcgcc tgaccattaa cggccatagc 420  
tattttaacc tggtgctggt gagcaacgtg ggcggcagcg gcgatgtgca tgcggtgtgg 480  
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cagagcaaca gctatctgga tggccagagc ctgagcttta ttgtgcgcgc gggcgatggc 600  
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<211> 224

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<213> Craterostigma plantagineum

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20 25 30

Thr Ala Ala Leu Ser Thr Thr Leu Phe Asn Asn Gly Leu Ala Cys Gly  
35 40 45

Ser Cys Tyr Gln Val Arg Cys Glu Gly Gly Pro Lys Trp Cys Val Arg  
50 55 60

Gly Gly Asp Arg Ile Ile Thr Val Thr Ala Thr Asn Phe Cys Pro Pro  
65 70 75 80

Asn Tyr Ala Leu Ala Asn Asp Asn Gly Gly Trp Cys Asn Pro Pro Arg  
85 90 95

Gln His Phe Asp Met Ala Gln Pro Ala Phe Val Arg Ile Ala His Tyr  
100 105 110

Arg Ala Gly Ile Val Pro Ile Ser Tyr Arg Arg Val Ser Cys Val Lys  
115 120 125

Lys Gly Gly Ile Arg Leu Thr Ile Asn Gly His Ser Tyr Phe Asn Leu  
130 135 140

Val Leu Val Ser Asn Val Gly Gly Ser Gly Asp Val His Ala Val Trp  
145 150 155 160

Ile Lys Gly Ser Gly Gly Gly Pro Trp Gln Ala Met Thr Arg Asn Trp  
165 170 175

Gly Gln Asn Trp Gln Ser Asn Ser Tyr Leu Asp Gly Gln Ser Leu Ser  
180 185 190

Phe Ile Val Arg Ala Gly Asp Gly Arg Thr Val Thr Ala Asn Asp Ile  
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Val Pro Arg Gly Trp Gln Phe Gly Gln Thr Phe Glu Gly Pro Gln Phe  
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sequence listing.txt

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sequence listing.txt

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sequence listing.txt

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